

Nucleic Acids Abstracts

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Annual Subject and Author

INDEXES

Volume 7 1977

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Nucleic Acids Abstracts

Annual Subject and Author

INDEXES

to Volume 7 1977

Editor Angela Pritchard

Assistant editor Eva Bornemann



ISSN 0048-1041

Compiled by
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Airfreight and mailing in the USA by
Publications Expediting Inc., Elmont, L.I.
200 Meacham Avenue, NY 11003

Printed by
Information Printing Ltd. Oxford, England

Published by Information Retrieval Ltd.,
London, England

Subject Index

Preface

From 1976, IRL has adopted a new system for the preparation of annual subject indexes. The system is based on the use of key word listings derived from the previous free-term indexes.

For each abstract, the indexer selects terms from a fixed list of key words which is continually updated.

The word list contains terms which will appear as entry points in the index, and subsidiary descriptor terms (e.g. aminoacylation, primary structure, purification).

From the terms selected, several chains of key words are derived, their word order being determined by the indexer.

The terms are separated by the following punctuation symbols:

1. comma - used to separate terms as in a normal sentence.
2. hyphen - used to indicate a relationship between terms.
3. oblique - used to indicate an alternative.
4. plus sign - used to indicate a combination
5. cross sign - used as an indication of hybrid formation.

Throughout the indexes cross-references are included. These are of one of the following types.

1. Directing the user from one synonym to another e.g. F-factor: see also colicinogenic factor, episome, plasmid, R-factor.
2. Guiding the user from the general to the more specific e.g. Antidiuretic hormone: see Vasopressin. .

WORKED EXAMPLE

Title of abstract: Characterization of the non-histone nuclear proteins associated with rapidly labelled heterogeneous nuclear RNA.

Entry terms: RNA (heterogeneous nuclear), Nonhistone protein.

Subsidiary descriptors: associated, characterization.

Index chains: a) RNA (heterogeneous nuclear), associated, nonhistone protein, characterization.
b) Nonhistone protein, RNA (heterogeneous nuclear) - associated, characterization.

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